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1644

RAW SEQUENCE LISTING

DATE: 02/19/2002

PATENT APPLICATION: US/09/847,208B

TIME: 12:13:27

Input Set : A:\UC067.002A-SEQ-ID.txt

Output Set: N:\CRF3\02192002\I847208B.raw

4 <110> APPLICANT: Saxon, Andrew
 5 Zhang, Ke
 6 Zhu, Daocheng
 8 <120> TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF
 9 IgE-MEDIATED ALLERGIC DISEASES
 12 <130> FILE REFERENCE: UC067.002A
 14 <140> CURRENT APPLICATION NUMBER: US 09/847,208B
 15 <141> CURRENT FILING DATE: 2001-05-01
 17 <160> NUMBER OF SEQ ID NOS: 177
 19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 696
 23 <212> TYPE: DNA
 24 <213> ORGANISM: Homo sapiens
 26 <400> SEQUENCE: 1
 27 gagcccaaat cttgtgacaa aactcacaca tgccccaccgt gccagcacc tgaactcctg 60
 28 gggggaccgt cagtcttctt cttcccccca aaaccaag acaccctcat gatctcccgg 120
 29 acccctgagg tcacatgcgt ggtggtggac gtgagccacg aagaccctga ggtcaagttc 180
 30 aactggtacg tggacggcgt ggaggtgcat aatgttaaga caaagccgcg ggaggagcag 240
 31 tacaacagca cgtaccgtgt ggtcagcgtc ctcaccgtcc tgcaccagaa ctggatgaat 300
 32 ggaaaggagt acaagtgc aaagtgc gcccagagaa ccacaggtgt acaccctgcc cccatcccgg 360
 33 atctccaaag ccaagaagca ggtcagcgtg acctgcctgg tcaaaggctt ctatcccagc 420
 34 gatgagctga ccaagaacca ggtcagcgtg acctgcctgg tcaaaggctt ctatcccagc 480
 35 gacatgcgcg tggagtggga gagcaatggg cagccggaga acaactacaa gaccacgcct 540
 36 cccgtgctgg actccgtcgg ctccttcttc ctctacagca agctcaccgt ggacaagagc 600
 37 aggtggcagc aggggaacgt cttctcatgc tccgtgatgc atgaggctct gcacaaccac 660
 38 taccagcaga ggagcctctc cctgtctccg ggtaaa 696
 40 <210> SEQ ID NO: 2
 41 <211> LENGTH: 330
 42 <212> TYPE: PRT
 43 <213> ORGANISM: Homo sapiens
 45 <400> SEQUENCE: 2
 46 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
 47 1 5 10 15
 48 Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
 49 20 25 30
 50 Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
 51 35 40 45
 52 Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser
 53 50 55 60
 54 Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
 55 65 70 75 80
 56 Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys

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57          85          90          95
58 Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
59          100          105          110
60 Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
61          115          120          125
62 Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
63          130          135          140
64 Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
65 145          150          155          160
66 Tyr Val Asp Gly Val Glu Val His Asn Val Lys Thr Lys Pro Arg Glu
67          165          170          175
68 Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
69          180          185          190
70 His Gln Asn Trp Met Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
71          195          200          205
72 Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Val
73          210          215          220
74 Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu
75 225          230          235          240
76 Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
77          245          250          255
78 Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
79          260          265          270
80 Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Val Gly Ser Phe Phe
81          275          280          285
82 Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
83          290          295          300
84 Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Gln
85 305          310          315          320
86 Gln Arg Ser Leu Ser Leu Ser Pro Gly Lys
87          325          330
90 <210> SEQ ID NO: 3
91 <211> LENGTH: 232
92 <212> TYPE: PRT
93 <213> ORGANISM: Homo sapiens
95 <400> SEQUENCE: 3
96 Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala
97 1          5          10          15
98 Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro
99          20          25          30
100 Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val
101          35          40          45
102 Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val
103          50          55          60
104 Asp Gly Val Glu Val His Asn Val Lys Thr Lys Pro Arg Glu Glu Gln
105 65          70          75          80
106 Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln
107          85          90          95
108 Asn Trp Met Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala

```

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```

109          100          105          110
110 Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Val Gln Pro
111          115          120          125
112 Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr
113          130          135          140
114 Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser
115 145          150          155          160
116 Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr
117          165          170          175
118 Lys Thr Thr Pro Pro Val Leu Asp Ser Val Gly Ser Phe Phe Leu Tyr
119          180          185          190
120 Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe
121          195          200          205
122 Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Gln Gln Arg
123          210          215          220
124 Ser Leu Ser Leu Ser Pro Gly Lys
125 225          230
128 <210> SEQ ID NO: 4
129 <211> LENGTH: 1445
130 <212> TYPE: DNA
131 <213> ORGANISM: Homo sapiens
133 <400> SEQUENCE: 4
134 tccacacaga gccatccgt cttccccttg acccgctgct gcaaaaacat tccctccaat 60
135 gccacctcgg tgactctggg ctgcctggcc acgggctact tcccggagcc ggtgatggtg 120
136 acctgggaca caggctccct caacgggaca actatgacct taccagccac caccctcacg 180
137 ctctctgggtc actatgccac catcagctct ctgacctctc cgggtgcgtg ggccaagcag 240
138 atgttcacct gccgtgtggc acacactcca tctgccacag actgggtoga caacaaaacc 300
139 ttcagcgtct gctccaggga cttcaccccg cccaccgtga agatcttaca gtcgtcctgc 360
140 gacggcgggcg ggcacttccc cccgaccatc cagctcctgt gcctcgtctc tgggtacacc 420
141 ccagggaacta tcaacatcac ctggctggag gacgggcagg tcatggacgt ggacttgtcc 480
142 accgcctcta ccacgcagga gggtagctg gcctccacac aaagcgagct caccctcagc 540
143 cagaagcact ggctgtcaga ccgcacctac acctgccagg tcacctatca aggtcacacc 600
144 tttgaggaca gcaccaagaa gtgtgcagat tccaaccoga gaggggtgag cgcctaccta 660
145 agccggccca gcccgttcga cctgttcatc cgcaagtcgc ccacgatcac ctgtctggtg 720
146 gtggacctgg caccagcaa ggggaccgtg aacctgacct ggtcccgggc cagtgggaag 780
147 cctgtgaacc actccaccag aaaggaggag aagcagcgca atggcacgtt aaccgtcacg 840
148 tccacctgc cgggtggcac ccgagactgg atcgaggggg agacctacca gtgcaggggtg 900
149 accaccccc acctgcccag ggccctcatg cgggtccacga ccaagaccag cggcccgcgt 960
150 gctgcccccg aagtctatgc gtttgcgacg ccggagtggc cggggagccg ggacaagcgc 1020
151 accctgcct gcctgatcca gaacttcatg cctgaggaca tctcgggtgca gtggctgcac 1080
152 aacgaggtgc agtcccga cgcccggcac agcacgacgc agccccgcaa gaccaagggc 1140
153 tccggttct togtcttcag ccgcctggag gtgaccaggg ccgaatggga gcagaaagat 1200
154 gagttcatct gccgtgcagt ccatgaggca gcgagcccct cacagaccgt ccagcgagcg 1260
155 gtgtctgtaa atcccggtaa atgacgtact cctgcctccc tccctcccag ggctccatcc 1320
156 agctgtgcag tggggaggac tggccagacc ttctgtccac tgttgcaatg accccaggaa 1380
157 gctaccccc ataaactgtg cctgctcaga gccccagtac acccattctt gggagcgggc 1440
158 agggc
160 <210> SEQ ID NO: 5
161 <211> LENGTH: 427

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162 <212> TYPE: PRT
163 <213> ORGANISM: Homo sapiens
165 <400> SEQUENCE: 5
166 Ser Thr Gln Ser Pro Ser Val Phe Pro Leu Thr Arg Cys Cys Lys Asn
167 1 5 10 15
168 Ile Pro Ser Asn Ala Thr Ser Val Thr Leu Gly Cys Leu Ala Thr Gly
169 20 25 30
170 Tyr Phe Pro Glu Pro Val Met Val Thr Trp Asp Thr Gly Ser Leu Asn
171 35 40 45
172 Gly Thr Thr Met Thr Leu Pro Ala Thr Thr Leu Thr Leu Ser Gly His
173 50 55 60
174 Tyr Ala Thr Ile Ser Leu Leu Thr Val Ser Gly Ala Trp Ala Lys Gln
175 65 70 75 80
176 Met Phe Thr Cys Arg Val Ala His Thr Pro Ser Ser Thr Asp Trp Val
177 85 90 95
178 Asp Asn Lys Thr Phe Ser Val Cys Ser Arg Asp Phe Thr Pro Pro Thr
179 100 105 110
180 Val Lys Ile Leu Gln Ser Ser Cys Asp Gly Gly Gly His Phe Pro Pro
181 115 120 125
182 Thr Ile Gln Leu Leu Cys Leu Val Ser Gly Tyr Thr Pro Gly Thr Ile
183 130 135 140
184 Asn Ile Thr Trp Leu Glu Asp Gly Gln Val Met Asp Val Asp Leu Ser
185 145 150 155 160
186 Thr Ala Ser Thr Thr Gln Glu Gly Glu Leu Ala Ser Thr Gln Ser Glu
187 165 170 175
188 Leu Thr Leu Ser Gln Lys His Trp Leu Ser Asp Arg Thr Tyr Thr Cys
189 180 185 190
190 Gln Val Thr Tyr Gln Gly His Thr Phe Glu Asp Ser Thr Lys Lys Cys
191 195 200 205
192 Ala Asp Ser Asn Pro Arg Gly Val Ser Ala Tyr Leu Ser Arg Pro Ser
193 210 215 220
194 Pro Phe Asp Leu Phe Ile Arg Lys Ser Pro Thr Ile Thr Cys Leu Val
195 225 230 235 240
196 Val Asp Leu Ala Pro Ser Lys Gly Thr Val Asn Leu Thr Trp Ser Arg
197 245 250 255
198 Ala Ser Gly Lys Pro Val Asn His Ser Thr Arg Lys Glu Glu Lys Gln
199 260 265 270
200 Arg Asn Gly Thr Leu Thr Val Thr Ser Thr Leu Pro Val Gly Thr Arg
201 275 280 285
202 Asp Trp Ile Glu Gly Glu Thr Tyr Gln Cys Arg Val Thr His Pro His
203 290 295 300
204 Leu Pro Arg Ala Leu Met Arg Ser Thr Thr Lys Thr Ser Gly Pro Arg
205 305 310 315 320
206 Ala Ala Pro Glu Val Tyr Ala Phe Ala Thr Pro Glu Trp Pro Gly Ser
207 325 330 335
208 Arg Asp Lys Arg Thr Leu Ala Cys Leu Ile Gln Asn Phe Met Pro Glu
209 340 345 350
210 Asp Ile Ser Val Gln Trp Leu His Asn Glu Val Gln Leu Pro Asp Ala
211 355 360 365

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```

212 Arg His Ser Thr Thr Gln Pro Arg Lys Thr Lys Gly Ser Gly Phe Phe
213      370      375      380
214 Val Phe Ser Arg Leu Glu Val Thr Arg Ala Glu Trp Glu Gln Lys Asp
215 385      390      395      400
216 Glu Phe Ile Cys Arg Ala Val His Glu Ala Ala Ser Pro Ser Gln Thr
217      405      410      415
218 Val Gln Arg Ala Val Ser Val Asn Pro Gly Lys
219      420      425
222 <210> SEQ ID NO: 6
223 <211> LENGTH: 320
224 <212> TYPE: PRT
225 <213> ORGANISM: Homo sapiens
227 <400> SEQUENCE: 6
228 Phe Thr Pro Pro Thr Val Lys Ile Leu Gln Ser Ser Cys Asp Gly Gly
229 1      5      10      15
230 Gly His Phe Pro Pro Thr Ile Gln Leu Leu Cys Leu Val Ser Gly Tyr
231      20      25      30
232 Thr Pro Gly Thr Ile Asn Ile Thr Trp Leu Glu Asp Gly Gln Val Met
233      35      40      45
234 Asp Val Asp Leu Ser Thr Ala Ser Thr Thr Gln Glu Gly Glu Leu Ala
235      50      55      60
236 Ser Thr Gln Ser Glu Leu Thr Leu Ser Gln Lys His Trp Leu Ser Asp
237 65      70      75      80
238 Arg Thr Tyr Thr Cys Gln Val Thr Tyr Gln Gly His Thr Phe Glu Asp
239      85      90      95
240 Ser Thr Lys Lys Cys Ala Asp Ser Asn Pro Arg Gly Val Ser Ala Tyr
241      100     105     110
242 Leu Ser Arg Pro Ser Pro Phe Asp Leu Phe Ile Arg Lys Ser Pro Thr
243      115     120     125
244 Ile Thr Cys Leu Val Val Asp Leu Ala Pro Ser Lys Gly Thr Val Asn
245      130     135     140
246 Leu Thr Trp Ser Arg Ala Ser Gly Lys Pro Val Asn His Ser Thr Arg
247 145     150     155     160
248 Lys Glu Glu Lys Gln Arg Asn Gly Thr Leu Thr Val Thr Ser Thr Leu
249      165     170     175
250 Pro Val Gly Thr Arg Asp Trp Ile Glu Gly Glu Thr Tyr Gln Cys Arg
251      180     185     190
252 Val Thr His Pro His Leu Pro Arg Ala Leu Met Arg Ser Thr Thr Lys
253      195     200     205
254 Thr Ser Gly Pro Arg Ala Ala Pro Glu Val Tyr Ala Phe Ala Thr Pro
255      210     215     220
256 Glu Trp Pro Gly Ser Arg Asp Lys Arg Thr Leu Ala Cys Leu Ile Gln
257 225     230     235     240
258 Asn Phe Met Pro Glu Asp Ile Ser Val Gln Trp Leu His Asn Glu Val
259      245     250     255
260 Gln Leu Pro Asp Ala Arg His Ser Thr Thr Gln Pro Arg Lys Thr Lys
261      260     265     270
262 Gly Ser Gly Phe Phe Val Phe Ser Arg Leu Glu Val Thr Arg Ala Glu
263      275     280     285

```

Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

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Input Set : A:\UC067.002A-SEQ-ID.txt

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L:3091 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82
L:3134 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L:3471 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93
L:3473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93
L:3488 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94
L:4523 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:126
L:6231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176
L:6233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176
L:6248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177
L:6250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177